

Resource Summary Report

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BSA4Yeast

RRID:SCR_017113

Type: Tool

Proper Citation

BSA4Yeast (RRID:SCR_017113)

Resource Information

URL: <https://bsa4yeast.lcsb.uni.lu>

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Description: Web application for Quantitative Trait Loci mapping via bulk segregant analysis of yeast sequencing data. Application provides automated data processing, annotations, and web interface to explore identified QTLs.

Synonyms: Bulk Segregant Analysis for Yeast

Resource Type: analysis service resource, data access protocol, data analysis service, production service resource, service resource, software resource, web service

Keywords: quantitative, trait, loci, mapping, bulk, segregant, analysis, yeast, sequencing, data, processing, annotation, bio.tools

Funding: Fonds Nationale de la Recherche (FNR) Luxembourg

Availability: Free, Freely available, Registration required

Resource Name: BSA4Yeast

Resource ID: SCR_017113

Alternate IDs: biotools:BSA4Yeast

Alternate URLs: <https://bio.tools/BSA4Yeast>

Record Creation Time: 20220129T080333+0000

Record Last Update: 20260919T195334+0000

Ratings and Alerts

No rating or validation information has been found for BSA4Yeast.

No alerts have been found for BSA4Yeast.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Zhang Z, et al. (2019) BSA4Yeast: Web-based quantitative trait locus linkage analysis and bulk segregant analysis of yeast sequencing data. GigaScience, 8(6).